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Lok, Si
Jaspers, Stephen R.

<130> 00-18

<151> 2000-03-10

<170> FastSEQ for Windows Version 3.0

<213> Homo sapiens

<222> (1)...(429)

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ggg gag ctg tgg ccg gga gct gag gcc cgg gca gcg cct tac ggg gtc 96
Gly Glu Leu Trp Pro Gly Ala Glu Ala Arg Ala Ala Pro Tyr Gly Val
20 25 30

agg ctt tgc ggc cga gaa ttc atc cga gca gtc atc ttc acc tgc ggg 144
Arg Leu Cys Gly Arg Glu Phe Ile Arg Ala Val Ile Phe Thr Cys Gly
35 40 45

ggc tcc cgg tgg aga cga tca gac atc ctg gcc cac gag gct atg gga 192
Gly Ser Arg Trp Arg Arg Ser Asp Ile Leu Ala His Glu Ala Met Gly
50 55 60

Met	Ala	Arg	Tyr	Met	Leu	Leu	Leu	Leu	Leu	Ala	Val	Trp	Val	Leu	Thr
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Gly	Glu	Leu	Trp	Pro	Gly	Ala	Glu	Ala	Arg	Ala	Ala	Pro	Tyr	Gly	Val
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Asp	Thr	Phe	Pro	Asp	Ala	Asp	Ala	Asp	Glu	Asp	Ser	Leu	Ala	Gly	Glu
65					70					75				80	
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<223> Cysteine motif

<221> VARIANT
<222> (3)...(13)
<223> Each Xaa is independently any amino acid residue
except cysteine.

<210>	4
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<221> VARIANT
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<223> Each Xaa is independently any amino acid residue
      except cysteine.
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mgngcngtna	thttyacntg	yggnggnwsn	mgntggmgnm	gnwsngayat	hytngcncay	180
gargcnatgg	gngayacntt	yccngaygcn	gaygcngayg	argaywsnyt	ngcnggngar	240
ytngaygarg	cnatgggnws	nwsngartgg	ytngcnytna	cnaarwsncc	ncargcntty	300
taymgnggnm	gnccnwsntg	gcarggnacn	ccnggngtny	tnmgnggnws	nmgngaygtn	360
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ytntgy						426

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<212>. DNA

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52